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=> s 9 -18:0-ACP desaturase
L1 24 9 -18:0-ACP DESATURASE

```
=> dup rem l1
PROCESSING COMPLETED FOR L1
L2          10 DUP REM L1 (14 DUPLICATES REMOVED)
```

=> s 12 and (mutant? or mutation? or variant?)
L3 8 L2 AND (MUTANT? OR MUTATION? OR VARIANT?)

=> s 13 and (114 or 117 or 118 or 179 or 181 or 188)
L4 2 L3 AND (114 OR 117 OR 118 OR 179 OR 181 OR 188)

=> d 14 1-8 ibib ab

L4 ANSWER 1 OF 2 CAPIUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 2000:881180 CAPIUS
DOCUMENT NUMBER: 134:52227
TITLE: General methods for directed mutagenesis and its application for **mutant** fatty acid desaturases
INVENTOR(S): Shanklin, John
PATENT ASSIGNEE(S): Brookhaven Science Associates, Llc, USA
SOURCE: PCT Int. Appl., 53 pp.
CODEN: PIXXD2
DOCUMENT TYPE: Patent
LANGUAGE: English
FAMILY ACC. NUM. COUNT: 3
PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2000075170	A1	20001214	WO 2000-US15741	20000608
W:	AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM			

RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY,
DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ,
CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
EP 1214334 A1 20020619 EP 2000-939675 20000608
R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
IE, SI, LT, LV, FI, RO, MK, CY, AL
PRIORITY APPLN. INFO.: US 1999-328550 A 19990609
WO 2000-US15741 W 20000608

AB The present invention relates to a method for producing **mutants** of a fatty acid desaturase, the unmutagenized precursor having 18-carbon chain substrate specificity, the produced **mutants** having substantially increased activity towards fatty acid substrates with chains contg. fewer than 18 carbons, relative to the activity of unmutagenized precursor desaturase. **Mutations** in the coding region of castor .**DELTA.9-18:0-ACP** desaturase which result in amino acid substitutions at position 114, 117, 118, 179, 181, or 188, and combined substitutions at positions 114 and 188, are described. A preferred bacterial strain used in the selection system, Escherichia coli MH13, is an unsatd. fatty auxotroph requiring exogenous unsatd. fatty acids to proliferate. Altered fatty acid compns. are demonstrated in transgenic Escherichia coli and Arabidopsis thaliana. DNA encoding the **mutants**, host cells contg. the DNA, transgenic plants expressing the DNA, and the desaturase enzyme produced by the method are also recited. Combinatorial full positional randomization of codons can be accomplished by a variety of methods. One such method is the use of overlap-extension PCR to replace all codons for candidate position amino acids with NNK or NNN. The process of overlap-extension PCR has been used to simultaneously introduce at least 9 independent **mutations** into a particular coding sequence. In another embodiment, a subset of one or more of the candidate positions are incompletely randomized, while the other candidate positions are fully randomized.

REFERENCE COUNT: 2 THERE ARE 2 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L4 ANSWER 2 OF 2 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 1998:395727 CAPLUS
DOCUMENT NUMBER: 129:146255
TITLE: A determinant of substrate specificity predicted from the acyl-acyl carrier protein desaturase of developing cat's claw seed
AUTHOR(S): Cahoon, Edgar B.; Shah, Salehuzzaman; Shanklin, John; Browse, John
CORPORATE SOURCE: Biology Department, Brookhaven National Laboratory, Upton, NY, 11976, USA
SOURCE: Plant Physiology (1998), 117(2), 593-598
CODEN: PLPHAY; ISSN: 0032-0889
PUBLISHER: American Society of Plant Physiologists
DOCUMENT TYPE: Journal
LANGUAGE: English

AB Cat's claw (*Doxantha unguis-cati* L.) vine accumulates nearly 80% palmitoleic acid (16:1.**DELTA.9**) plus cis-vaccenic acid (18:1.**DELTA.11**) in its seed oil. To characterize the biosynthetic origin of these unusual fatty acids, cDNAs for acyl-acyl carrier protein (acyl-ACP) desaturases were isolated from developing cat's claw seeds. The predominant acyl-ACP desaturase cDNA identified encoded a polypeptide that is closely related to the stearoyl (.**DELTA.9-18:0-ACP** desaturase from castor (*Ricinus communis* L.) and other species. Upon expression in Escherichia coli, the cat's claw polypeptide functioned as a .**DELTA.9** acyl-ACP desaturase but displayed a distinct substrate specificity for palmitate (16:0)-ACP rather than stearate (18:0)-ACP. Comparison of the predicted amino acid sequence of the cat's claw enzyme

with that of the castor .DELTA.9-18:0-ACP desaturase suggested that a single amino acid substitution (L118W) might account in large part for the differences in substrate specificity between the two desaturases. Consistent with this prediction, conversion of leucine-118 to tryptophan in the mature castor .DELTA.9-18:0-ACP

desaturase resulted in an 80-fold increase in the relative specificity of this enzyme for 16:0-ACP. The alteration in substrate specificity obsd. in the L118W mutant is in agreement with a crystallog. model of the proposed substrate-binding pocket of the castor .DELTA.9-18:0-ACP

desaturase.

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PRIORITY APPLN. INFO.:			R: AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT, IE, SI, LT, LV, FI, RO, MK, CY, AL	US 1999-328550 A 19990609
				WO 2000-US15741 W 20000608

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acid compns. are demonstrated in transgenic Escherichia coli and Arabidopsis thaliana. DNA encoding the **mutants**, host cells contg. the DNA, transgenic plants expressing the DNA, and the desaturase enzyme produced by the method are also recited. Combinatorial full positional randomization of codons can be accomplished by a variety of methods. One such method is the use of overlap-extension PCR to replace all codons for candidate position amino acids with NNK or NNN. The process of overlap-extension PCR has been used to simultaneously introduce at least 9 independent **mutations** into a particular coding sequence. In another embodiment, a subset of one or more of the candidate positions are incompletely randomized, while the other candidate positions are fully randomized.

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DOCUMENT NUMBER: 129:146255
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AUTHOR(S): Cahoon, Edgar B.; Shah, Salehuzzaman; Shanklin, John; Browse, John
CORPORATE SOURCE: Biology Department, Brookhaven National Laboratory, Upton, NY, 11976, USA
SOURCE: Plant Physiology (1998), 117(2), 593-598
CODEN: PLPHAY; ISSN: 0032-0889
PUBLISHER: American Society of Plant Physiologists
DOCUMENT TYPE: Journal
LANGUAGE: English

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REFERENCE COUNT: 20 THERE ARE 20 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

=> d 14 3-8 ibib ab

2 ANSWERS ARE AVAILABLE. SPECIFIED ANSWER NUMBER EXCEEDS ANSWER SET SIZE
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ENTER ANSWER NUMBER OR RANGE (1):d 14 3 ibib ab
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several answer numbers, Example: 3,7,10
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or a combination of these. Example: 3,7,9-10,15
ENTER ANSWER NUMBER OR RANGE (1):end

=> d his

(FILE 'HOME' ENTERED AT 11:13:45 ON 19 MAR 2003)

FILE 'MEDLINE, CAPLUS, BIOSIS, BIOTECHDS, SCISEARCH, EMBASE' ENTERED AT
11:14:30 ON 19 MAR 2003

L1 24 S 9 -18:0-ACP DESATURASE
L2 10 DUP REM L1 (14 DUPLICATES REMOVED)
L3 8 S L2 AND (MUTANT? OR MUTATION? OR VARIANT?)
L4 2 S L3 AND (114 OR 117 OR 118 OR 179 OR 181 OR 188)

=> d 13 1-8 ibib ab

L3 ANSWER 1 OF 8 MEDLINE
ACCESSION NUMBER: 2001331977 MEDLINE
DOCUMENT NUMBER: 21293111 PubMed ID: 11294879
TITLE: Engineering delta 9-16:0-acyl carrier protein (ACP)
desaturase specificity based on combinatorial saturation
mutagenesis and logical redesign of the castor delta
9-18:0-ACP
desaturase.
AUTHOR: Whittle E; Shanklin J
CORPORATE SOURCE: Biology Department, Brookhaven National Laboratory, Upton,
New York 11973, USA.
SOURCE: JOURNAL OF BIOLOGICAL CHEMISTRY, (2001 Jun 15) 276 (24)
21500-5.
Journal code: 2985121R. ISSN: 0021-9258.
PUB. COUNTRY: United States
DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)
LANGUAGE: English
FILE SEGMENT: Priority Journals
ENTRY MONTH: 200107
ENTRY DATE: Entered STN: 20010723
Last Updated on STN: 20030105
Entered Medline: 20010719

AB Six amino acid locations in the soluble castor Delta(9)-18:0-acyl carrier protein (ACP) desaturase were identified that can affect substrate specificity. Combinatorial saturation mutagenesis of these six amino acids, in conjunction with selection, using an unsaturated fatty acid auxotroph system, led to the isolation of variants with up to 15-fold increased specific activity toward 16-carbon substrates. The most improved mutant, com2, contained two substitutions (T117R/G188L) common to five of the 19 complementing variants subjected to further analysis. These changes, when engineered into otherwise wild-type 18:0-ACP desaturase to make mutant 5.2, produced a 35-fold increase in specific activity with respect to 16-carbon substrates. Kinetic analysis revealed changes in both k(cat) and K(m) that result in an 82-fold improvement in specificity factor for 16-carbon substrate compared with wild-type enzyme. Improved substrate orientation apparently compensated for loss of binding energy that results from the loss of desolvation energy for 16-carbon substrates. Mutant 5.2 had specific activity for 16-carbon substrates 2 orders of magnitude higher than those of known natural 16-carbon specific desaturases. These data support the hypothesis that it should be possible to reengineer archetypal enzymes to achieve substrate specificities characteristic of recently evolved enzymes while retaining the desired stability and/or turnover characteristics of a parental paralog.

L3 ANSWER 2 OF 8 MEDLINE
ACCESSION NUMBER: 2001039056 MEDLINE
DOCUMENT NUMBER: 20504514 PubMed ID: 11027301
TITLE: Substrate-dependent **mutant** complementation to select fatty acid desaturase **variants** for metabolic engineering of plant seed oils.
AUTHOR: Cahoon E B; Shanklin J
CORPORATE SOURCE: Biology Department, Brookhaven National Laboratory, Upton, NY 11973, USA.
SOURCE: PROCEEDINGS OF THE NATIONAL ACADEMY OF SCIENCES OF THE UNITED STATES OF AMERICA, (2000 Oct 24) 97 (22) 12350-5.
Journal code: 7505876. ISSN: 0027-8424.
PUB. COUNTRY: United States.
DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)
LANGUAGE: English
FILE SEGMENT: Priority Journals
ENTRY MONTH: 200011
ENTRY DATE: Entered STN: 20010322
Last Updated on STN: 20010322
Entered Medline: 20001128

AB We demonstrate that naturally occurring C(14) and C(16)-specific acyl-acyl carrier protein (ACP) desaturases from plants can complement the unsaturated fatty acid (UFA) auxotrophy of an *Escherichia coli* fabA/fadR **mutant**. Under the same growth conditions, C(18)-specific delta(9)-stearoyl (18:0)-ACP desaturases are unable to complement the UFA auxotrophy. This difference most likely results from the presence of sufficient substrate pools of C(14) and C(16) acyl-ACPs but a relative lack of C(18) acyl-ACP pools in *E. coli* to support the activities of the plant fatty acid desaturase. Based on this, a substrate-dependent selection system was devised with the use of the *E. coli* UFA auxotroph to isolate **mutants** of the castor delta(9)-18:
O-ACP desaturase that display enhanced specificity for C(14) and C(16) acyl-ACPs. Using this selection system, a number of desaturase **variants** with altered substrate specificities were isolated from pools of randomized **mutants**. These included several G188L **mutant** isolates, which displayed a 15-fold increase in specific activity with 16:0-ACP relative to the wild-type castor delta(9)-18:0-ACP **desaturase**. Expression of this **mutant** in *Arabidopsis thaliana* resulted in the accumulation of unusual monounsaturated fatty acids to amounts of >25% of the seed oil. The bacterial selection system described here thus provides a rapid means of isolating **variant** fatty acid desaturase activities for modification of seed oil composition.

L3 ANSWER 3 OF 8 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 2002:586274 CAPLUS
DOCUMENT NUMBER: 137:307376
TITLE: What limits production of unusual monoenoic fatty acids in transgenic plants?
AUTHOR(S): Suh, Mi Chung; Schultz, David J.; Ohlrogge, John B.
CORPORATE SOURCE: Department of Plant Biology, Michigan State University, East Lansing, MI, 48824, USA
SOURCE: Planta (2002), 215(4), 584-595
CODEN: PLANAB; ISSN: 0032-0935
PUBLISHER: Springer-Verlag
DOCUMENT TYPE: Journal
LANGUAGE: English
AB Unusual monounsaturated fatty acids are major constituents (greater than 80%) in seeds of *Coriandrum sativum* L. (coriander) and *Thunbergia alata* Bojer, as well as in glandular trichomes (greater than 80% derived products) of *Pelargonium .times.hortorum* (geranium). These diverged fatty acid structures are produced via distinct plastidial acyl-acyl carrier protein

(ACP) desaturases. When expressed in *Arabidopsis thaliana* (L.) Heynh. under strong seed-specific promoters the unusual acyl-ACP desaturases resulted in accumulation of unusual monoene fatty acids at 1-15% of seed fatty acid mass. In this study, we have examd. several factors that potentially limit higher prodn. of unusual monoenes in transgenic oilseeds. (i) Immunoblots indicated that the introduced desaturases were expressed at levels equiv. to or higher than the endogenous .DELTA.

9 18:0-ACP desaturase.

However, the level of unusual fatty acid produced in transgenic plants was not correlated with the level of desaturase expression. (ii) The unusual desaturases were expressed in several backgrounds, including antisense 18:0-ACP desaturase plants, in fab1 **mutants**, and co-expressed with specialized ACP or ferredoxin isoforms. None of these expts. led to high prodn. of expected products. (iii) No evidence was found for degrdn. of the unusual fatty acids during seed development. (iv) Petroselinic acid added to developing seeds was incorporated into triacylglycerol as readily as oleic acid, suggesting no major barriers to its metab. by enzymes of glycerolipid assembly. (v) In vitro and in situ assay of acyl-ACP desaturases revealed a large discrepancy of activity when comparing unusual acyl-ACP desaturases with the endogenous .DELTA.9

18:0-ACP desaturase. The combined results, coupled with the sensitivity of acyl-ACP desaturase activity to centrifugation and low salt or detergent suggests low prodn. of unusual monoenes in transgenic plants may be due to the lack of, or incorrect assemble of, a necessary multi-component enzyme assocn.

REFERENCE COUNT: 48 THERE ARE 48 CITED REFERENCES AVAILABLE FOR THIS RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L3 ANSWER 4 OF 8 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 2000:881180 CAPLUS
DOCUMENT NUMBER: 134:52227
TITLE: General methods for directed mutagenesis and its application for **mutant** fatty acid desaturases
INVENTOR(S): Shanklin, John
PATENT ASSIGNEE(S): Brookhaven Science Associates, Llc, USA
SOURCE: PCT Int. Appl., 53 pp.
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PATENT INFORMATION:

PATENT NO.	KIND	DATE	APPLICATION NO.	DATE
WO 2000075170	A1	20001214	WO 2000-US15741	20000608
W: AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM				
RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG				
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PRIORITY APPLN. INFO.:			US 1999-328550	A 19990609
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desaturase which result in amino acid substitutions at position 114, 117, 118, 179, 181, or 188, and combined substitutions at positions 114 and 188, are described. A preferred bacterial strain used in the selection system, Escherichia coli MH13, is an unsatd. fatty auxotroph requiring exogenous unsatd. fatty acids to proliferate. Altered fatty acid compns. are demonstrated in transgenic Escherichia coli and Arabidopsis thaliana. DNA encoding the **mutants**, host cells contg. the DNA, transgenic plants expressing the DNA, and the desaturase enzyme produced by the method are also recited. Combinatorial full positional randomization of codons can be accomplished by a variety of methods. One such method is the use of overlap-extension PCR to replace all codons for candidate position amino acids with NNK or NNN. The process of overlap-extension PCR has been used to simultaneously introduce at least 9 independent **mutations** into a particular coding sequence. In another embodiment, a subset of one or more of the candidate positions are incompletely randomized, while the other candidate positions are fully randomized.

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L3 ANSWER 5 OF 8 CAPLUS COPYRIGHT 2003 ACS

ACCESSION NUMBER: 2000:759826 CAPLUS

DOCUMENT NUMBER: 134:39564

TITLE: Stearoyl-acyl carrier protein and unusual acyl-acyl carrier protein desaturase activities are differentially influenced by ferredoxin

AUTHOR(S): Schultz, David J.; Suh, Mi Chung; Ohlrogge, John B.

CORPORATE SOURCE: Department of Botany and Plant Pathology, Michigan State University, East Lansing, MI, 48824, USA

SOURCE: Plant Physiology (2000), 124(2), 681-692

CODEN: PLPHAY; ISSN: 0032-0889

PUBLISHER: American Society of Plant Physiologists

DOCUMENT TYPE: Journal

LANGUAGE: English

AB Acyl-acyl carrier protein (ACP) desaturases function to position a single double bond into an acyl-ACP substrate and are best represented by the ubiquitous .DELTA.9 18:0-ACP

desaturase. Several **variant** acyl-ACP desaturases have also been identified from species that produce unusual monoenoic fatty acids. All known acyl-ACP desaturase enzymes use ferredoxin as the electron-donating cofactor, and in almost all previous studies the photosynthetic form of ferredoxin rather than the non-photosynthetic form has been used to assess activity. The influence of different forms of ferredoxin on acyl-ACP desaturases was examd. Using combinations of in vitro acyl-ACP desaturase assays and [14C]malonyl-CoA labeling studies, it was detd. that heterotrophic ferredoxin isoforms support up to 20-fold higher unusual acyl-ACP desaturase activity in coriander (*Coriandrum sativum*), *Thunbergia alata*, and garden geranium (*Pelargonium .times. hortorum*) when compared with photosynthetic ferredoxin isoforms. Heterotrophic ferredoxin also increases activity of the ubiquitous .DELTA.9 18:0-ACP **desaturase**

1.5-3.0-fold in both seed and leaf exts. These results suggest that ferredoxin isoforms may specifically interact with acyl-ACP desaturases to achieve optimal enzyme activity and that heterotrophic isoforms of ferredoxin may be the in vivo electron donor for this reaction.

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L3 ANSWER 6 OF 8 CAPLUS COPYRIGHT 2003 ACS

ACCESSION NUMBER: 1998:395727 CAPLUS
DOCUMENT NUMBER: 129:146255
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L3 ANSWER 7 OF 8 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 1997:333116 CAPLUS
DOCUMENT NUMBER: 127:46988
TITLE: Redesign of soluble fatty acid desaturases from plants from altered substrate specificity and double bond position
AUTHOR(S): Cahoon, Edgar B.; Lindqvist, Ylva; Schneider, Gunter; Shanklin, John
CORPORATE SOURCE: Biol. Dep., Brookhaven Natl. Lab., Upton, NY, 11973, USA
SOURCE: Proceedings of the National Academy of Sciences of the United States of America (1997), 94(10), 4872-4877
CODEN: PNASA6; ISSN: 0027-8424
PUBLISHER: National Academy of Sciences
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AB Acyl-acyl carrier protein (ACP) desaturases introduce double bonds at specific positions in fatty acids of defined chain lengths and are one of the major determinants of the monounsatd. fatty acid compn. of vegetable oils. Mutagenesis studies were conducted to det. the structural basis for the substrate and double bond positional specificities displayed by acyl-ACP desaturases. By replacement of specific amino acid residues in a

.DELTA.6-palmitoyl (16:0)-ACP desaturase with their equiv. from a .DELTA.9-stearoyl (18:0)-ACP desaturase, **mutant** enzymes were identified that have altered fatty acid chain-length specificities or that can insert double bonds into either the .DELTA.6 or .DELTA.9 positions of 16:0- and 18:0-ACP. Most notably, by replacement of 5 amino acids (A181T/A200F/S205N/L206T/G207A), the .DELTA.6-16:0-AcP desaturase was converted into an enzyme that functions principally as a .DELTA.9-18:0-ACP desaturase. Many of the determinants of fatty acid chain-length specificity in these **mutants** are found in residues that line the substrate binding channel as revealed by x-ray crystallog. of the .DELTA.9-18:0-ACP desaturase. The crystallog. model of the active site is also consistent with the diverged activities assocd. with naturally occurring **variant** acyl-ACP desaturases. In addn., on the basis of the active-site model, a .DELTA.9-18:0-AcP desaturase was converted into an enzyme with substrate preference for 16:0-ACP by replacement of 2 residues (L118F/P179I). These results demonstrate the ability to rationally modify acyl-ACP desaturase activities through site-directed mutagenesis and represent a first step toward the design of acyl-AcP desaturases for the prodn. of novel monounsatd. fatty acids in transgenic oilseed crops.

L3 ANSWER 8 OF 8 CAPLUS COPYRIGHT 2003 ACS
ACCESSION NUMBER: 1997:330585 CAPLUS
DOCUMENT NUMBER: 127:46978
TITLE: Characterization of a structurally and functionally diverged acyl-acyl carrier protein desaturase from milkweed seed
AUTHOR(S): Cahoon, Edgar B.; Coughlan, Sean J.; Shanklin, John
CORPORATE SOURCE: Biology Department, Brookhaven National Laboratory, Upton, NY, 11973, USA
SOURCE: Plant Molecular Biology (1997), 33(6), 1105-1110
CODEN: PMBIDB; ISSN: 0167-4412
PUBLISHER: Kluwer
DOCUMENT TYPE: Journal
LANGUAGE: English
AB A cDNA for a structurally **variant** acyl-acyl carrier protein (ACP) desaturase was isolated from milkweed (*Asclepias syriaca*) seed, a tissue enriched in palmitoleic (16:1.DELTA.9) and cis-vaccenic (18:1.DELTA.11) acids. Exts. of *Escherichia coli* that express the milkweed cDNA catalyzed .DELTA.9 desatn. of acyl-ACP substrates, and the recombinant enzyme exhibited seven- to ten-fold greater specificity for palmitoyl (16:0)-ACP and 30-fold greater specificity for myristoyl (14:0)-ACP than did known .DELTA.9-stearoyl (18:0)-ACP desaturases. Like other **variant** acyl-ACP desaturases reported to date, the milkweed enzyme contains fewer amino acids near its N-terminus compared to previously characterized .DELTA.9-18:0-ACP desaturases. Based on the activity of an N-terminal deletion **mutant** of a .DELTA.9-18:0-ACP desaturase, this structural feature likely does not account for differences in substrate specificities.

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(FILE 'HOME' ENTERED AT 11:13:45 ON 19 MAR 2003)

FILE 'MEDLINE, CAPLUS, BIOSIS, BIOTECHDS, SCISEARCH, EMBASE' ENTERED AT 11:14:30 ON 19 MAR 2003

L1 24 S 9 -18:0-ACP DESATURASE
L2 10 DUP REM L1 (14 DUPLICATES REMOVED)
L3 8 S L2 AND (MUTANT? OR MUTATION? OR VARIANT?)

L4 2 S L3 AND (114 OR 117 OR 118 OR 179 OR 181 OR 188)

=> log y

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